## SEQUENCE LISTING

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<110> TERLECKY, Stanley R.
      WALTON, Paul A.
<120> PROMOTION OF PEROXISOMAL CATALASE FUNCTION IN CELLS
<130> 28928.0009)
<140> (to be assigned)
<141> 2005-04-29
<150> PCT/US03/34512
<151> 2003-10-30
<150> US 60/422,100
<151> 2002-10-30
<160> 21
<170> PatentIn Ver. 3.2
<210> 1
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Lys Ala Asn Leu
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<213> Homo sapiens
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Met Ala Asp Ser Arg Asp Pro Ala Ser Asp Gln Met Gln His Trp Lys
                  5
Glu Gln Arg Ala Ala Gln Lys Ala Asp Val Leu Thr Thr Gly Ala Gly
             20
Asn Pro Val Gly Asp Lys Leu Asn Val Ile Thr Val Gly Pro Arg Gly
                              40
Pro Leu Leu Val Gln Asp Val Val Phe Thr Asp Glu Met Ala His Phe
                         55
Asp Arg Glu Arg Ile Pro Glu Arg Val Wal His Ala Lys Gly Ala Gly
                     70
 65
                                          75
                                                               80
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Ala Phe Gly Tyr Phe Glu Val Thr His Asp Ile Thr Lys Tyr Ser Lys Ala Lys Val Phe Glu His Ile Gly Lys Lys Thr Pro Ile Ala Val Arg 105 Phe Ser Thr Val Ala Gly Glu Ser Gly Ser Ala Asp Thr Val Arg Asp Pro Arg Gly Phe Ala Val Lys Phe Tyr Thr Glu Asp Gly Asn Trp Asp Leu Val Gly Asn Asn Thr Pro Ile Phe Phe Ile Arg Asp Pro Ile Leu 145 150 Phe Pro Ser Phe Ile His Ser Gln Lys Arg Asn Pro Gln Thr His Leu Lys Asp Pro Asp Met Val Trp Asp Phe Trp Ser Leu Arg Pro Glu Ser 185 Leu His Gln Val Ser Phe Leu Phe Ser Asp Arg Gly Ile Pro Asp Gly His Arg His Met Asn Gly Tyr Gly Ser His Thr Phe Lys Leu Val Asn Ala Asn Gly Glu Ala Val Tyr Cys Lys Phe His Tyr Lys Thr Asp Gln Gly Ile Lys Asn Leu Ser Val Glu Asp Ala Ala Arg Leu Ser Gln Glu 250 Asp Pro Asp Tyr Gly Ile Arg Asp Leu Phe Asn Ala Ile Ala Thr Gly Lys Tyr Pro Ser Trp Thr Phe Tyr Ile Gln Val Met Thr Phe Asn Gln 275 280 Ala Glu Thr Phe Pro Phe Asn Pro Phe Asp Leu Thr Lys Val Trp Pro 295 His Lys Asp Tyr Pro Leu Ile Pro Val Gly Lys Leu Val Leu Asn Arg 310 315 Asn Pro Val Asn Tyr Phe Ala Glu Val Glu Gln Ile Ala Phe Asp Pro 325 335 Ser Asn Met Pro Pro Gly Ile Glu Ala Ser Pro Asp Lys Met Leu Gln 345 Gly Arg Leu Phe Ala Tyr Pro Asp Thr His Arg His Arg Leu Gly Pro 360 Asn Tyr Leu His Ile Pro Val Asn Cys Pro Tyr Arg Ala Arg Val Ala

375

380

370

Asn Tyr Gln Arg Asp Gly Pro Met Cys Met Gln Asp Asn Gln Gly Gly 390 Ala Pro Asn Tyr Tyr Pro Asn Ser Phe Gly Ala Pro Glu Gln Gln Pro 405 410 Ser Ala Leu Glu His Ser Ile Gln Tyr Ser Gly Glu Val Arg Arg Phe 420 Asn Thr Ala Asn Asp Asp Asn Val Thr Gln Val Arg Ala Phe Tyr Val 440 Asn Val Leu Asn Glu Glu Gln Arg Lys Arg Leu Cys Glu Asn Ile Ala 460 Gly His Leu Lys Asp Ala Gln Ile Phe Ile Gln Lys Lys Ala Val Lys 470 Asn Phe Thr Glu Val His Pro Asp Tyr Gly Ser His Ile Gln Ala Leu 490 Leu Asp Lys Tyr Asn Ala Glu Lys Pro Lys Asn Ala Ile His Thr Phe 500 505 Val Gln Ser Gly Ser His Leu Ala Ala Arg Glu Lys Ala Asn Leu 515 520 525 <210> 3 <211> 1586 <212> DNA <213> Homo sapiens <400> 3 atggctgaca gccgggatcc cgccagcgac cagatgcagc actggaagga gcagcgggcc 60 gcgcagaaag ctgatgtcct gaccactgga gctggtaacc cagtaggaga caaacttaat 120 gttattacag tagggccccg tgggcccctt cttgttcagg atgtggtttt cactgatgaa 180 atggctcatt ttgaccgaga gagaattcct gagagagttg tgcatgctaa aggagcaggg 240 gcctttggct actttgaggt cacacatgac attaccaaat actccaaggc aaaggtattt 300 gagcatattg gaaagaagac tcccatcgca gttcggttct ccactgttgc tggagaatcg 360 ggttcagctg acacagttcg ggaccctcgt gggtttgcag tgaaatttta cacagaagat 420 ggtaactggg atctcgttgg aaataacacc cccattttct tcatcaggga tcccatattg 480 tttccatctt ttatccacag ccaaaagaga aatcctcaga cacatctgaa ggatccggac 540 atggtctggg acttctggag cctacgtcct gagtctctgc atcaggtttc tttcttgttc 600 agtgatcggg ggattccaga tggatccatc gccacatgaa tggatatgga tcacatactt 660 tcaagctggt taatgcaaat ggggaggcag tttattgcaa attccattat aagactgacc 720 agggcatcaa aaacctttct gttgaagatg cggcgagact ttcccaggaa gatcctgact 780 atggcatccg ggatcttttt aacgccattg ccacaggaaa gtacccctcc tggacttttt 840 acatccaggt catgacattt aatcaggcag aaacttttcc atttaatcca ttcgatctca 900 ccaaggtttg gcctcacaag gactaccctc tcatcccagt tggtaaactg gtcttaaacc 960 ggaatccagt taattacttt gctgaggttg aacagatagc cttcgaccca agcaacatgc 1020 cacctggcat tgaggccagt cctgacaaaa tgcttcaggg ccgccttttt gcctatcctg 1080

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<223> Description of Artificial Sequence: Synthetic
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Arg Gln Ile Lys Ile Phe Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
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                                     10
<210> 6
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<212> PRT
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peptide

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<400> 7
Arg Gln Ile Lys Ile Phe Phe Gln Asn Arg Arg Met Lys Phe Lys Lys
                  5
<210> 8
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Lys Glu Thr Trp Trp Glu Thr Trp Trp Thr Glu Trp Ser Gln Pro Lys
Lys Lys Arg Lys Val
             20
<210> 9
<211> 12
<212> PRT
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     motif
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<210> 10
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<400> 10
Lys Lys Lys Arg Lys Val
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Arg Leu Gln Val Val Leu Gly His Leu
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<400> 12
gtgaaccgtc agatccgct
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<210> 13
<211> 50
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cgtctcgagt tatagatcag ctttcagctc gtccatgccg agagtgatcc
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<210> 14
<211> 30
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<210> 15
<211> 42
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cgtctcgagt tataatttgg acaggaacag gtggtggcgg cc
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<211> 31
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<400> 16
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<210> 17
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<210> 18
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